©02 Rec'd PCT/PTO 3 0 JAN 2001

SEQUENCE LISTING

<110> FROHBERG, Claus

<120> NUCLEIC ACID MOLECULES ENCODING AN ALPHA-GLUCOSIDASE, PLANTS WHICH SYNTHESIZE A MODIFIED STARCH, THE GENERATION OF THE PLANTS, THEIR USE, AND THE MODIFIED STARCH

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250

Thr Pro Ser Ser Thr Phe Asp Asp Pro Pro Tyr Lys Ile Asn Asn Ser

245

255



260 265 270

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Ile Leu Lys Gln Gly Ala Thr Ser Val Asp Ala Tyr Phe Pro Ala Gly

Asn Trp Phe Asp Leu Phe Asn Tyr Ser Arg Ser Val Ser Leu Asn Gln 500 505 510

Gly Thr Tyr Met Thr Leu Asp Ala Pro Pro Asp His Ile Asn Val His 515 520 525

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Ser Lys Asn Ser Thr Gly Glu Leu Phe Val Asp Asp Asp Glu Val
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Gln Met Gly Arg Glu Gly Gly Arg Trp Thr Leu Val Lys Phe Asn Ser 580 585 590

Asn Ile Ile Gly Asn Lys Ile Val Val Lys Ser Glu Val Val Asn Gly 595 600 605

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Gly Phe Glu Asn Val Arg Gly Leu Lys Ser Tyr Glu Leu Val Gly Ser 625 630 635 640

His Gln Gln Gly Asn Thr Thr Met Lys Glu Ser Leu Lys Gln Ser Gly 645 650 655

Gln Phe Val Thr Met Glu Ile Ser Gly Met Ser Ile Leu Ile Gly Lys 660 665 670

Glu Phe Lys Leu Glu Leu Tyr Ile Ile Thr 675 680

JC02 Rec'd PCT/PTO 3 0 JAN 2001

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SEQUENCE LISTING

<110> Hoechst Schering AgrEvo GmbH

Nucleic acid molecules encoding an \(\alpha\)-glucosidase, plants which synthesize a modified starch, the generation of the plants, their use, and the modified starch

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AGCTTCACCG	TCCGCCGGCG	CTCCACCGGG	GATACTCTTT	TCGATACTTC	GCCGGAGTTA	240
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R KR X

REPLACEMENT SHEET (RULE 26)

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Ile Val Tyr Thr Gly Asp Arg Ile Ser Tyr Lys Val Ile Gly Gly Leu

Ile Asp Leu Tyr Phe Phe Ala Gly Pro Ser Pro Glu Met Val Val Asp
65 70 75 80

Gln Tyr Thr Gln Leu Ile Gly Arg Pro Ala Ala Met Pro Tyr Trp Ser

Phe Gly Phe His Gln Cys Arg Trp Gly Tyr Lys Asn Ile Asp Asp Val

Glu Leu Val Val Asp Ser Tyr Ala Lys Ser Arg Ile Pro Leu Glu Val 115 120 125

Met Trp Thr Asp Ile Asp Tyr Met Asp Gly Phe Lys Asp Phe Thr Leu 130 135 140

Asp Pro Val Asn Phe Pro Leu Glu Arg Val Ile Phe Phe Leu Arg Lys 145 150 155 160

Leu His Gln Asn Asp Gln Lys Tyr Val Leu Ile Val Asp Pro Gly Ile 165 170 175

Ser Ile Asn Asn Thr Tyr Asp Thr Tyr Arg Arg Gly Met Glu Ala Asp 180 185 190

Val Phe Ile Lys Arg Asp Asn Met Pro Tyr Gln Gly Val Val Trp Pro 195 200 205

Gly Asn Val Tyr Tyr Pro Asp Phe Leu Asn Pro Ala Thr Glu Val Phe

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Thr Pro Ser Ser Thr Phe Asp Asp Pro Pro Tyr Lys Ile Asn Asn Ser 260 265 270

Gly Asp His Leu Pro Ile Asn Tyr Arg Thr Val Pro Ala Thr Ser Thr 275 280 285

His Phe Gly Asp Thr Met Glu Tyr Asn Val His Asn Leu Tyr Gly Leu 290 295 300

Leu Glu Ser Arg Ala Thr Tyr Ser Ala Leu Val Asn Val Thr Gly Lys 305 310 315 320

Arg Pro Phe Ile Leu Val Arg Ser Thr Phe Leu Gly Ser Gly Arg Tyr 325 330 335

Thr Ser His Trp Thr Gly Asp Asn Ala Ala Thr Trp Asn Asp Leu Ala 340 . 345 . 350

Tyr Ser Ile Pro Thr Ile Leu Ser Phe Gly Leu Phe Gly Ile Pro Met 355 360 365

Val Gly Ala Asp Ile Cys Gly Phe Ser Ser Asn Thr Thr Glu Glu Leu 370 375 380

Cys Arg Arg Trp Ile Gln Leu Gly Ala Phe Tyr Pro Phe Ala Arg Asp 385 390 395

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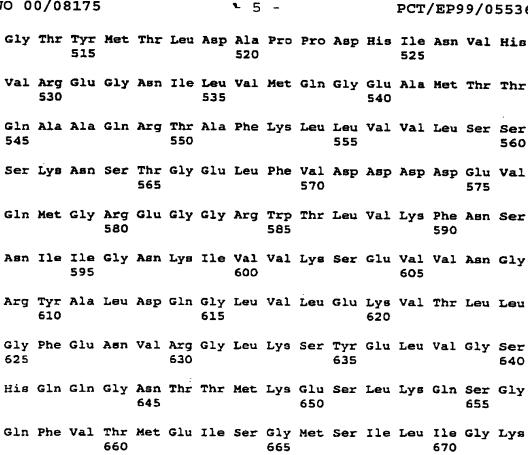
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Asn Trp Phe Asp Leu Phe Asn Tyr Ser Arg Ser Val Ser Leu Asn Gln 500 505



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